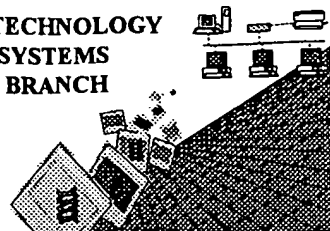


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BIOTECHNOLOGY
SYSTEMS
BRANCH0590
0930**RAW SEQUENCE LISTING**
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



OIPE

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/901,572

DATE: 10/01/2002

TIME: 16:12:31

Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

3 <110> APPLICANT: Nippon Zeon Co., Ltd.,
W--> 4 <120> TITLE OF INVENTION: Modified DNA molecules, Recombinants and uses thereof
W--> 5 <130> FILE REFERENCE: J209
W--> 6 <140> CURRENT APPLICATION NUMBER:
C--> 7 <141> CURRENT FILING DATE: 2002-09-23
E--> 8 <160> NUMBER OF SEQ ID NOS:
9 <170> SOFTWARE: PatentIn Ver. 2.1

mandatory response needed
only 4 sequences in submitted file

Does Not Comply
Corrected Diskette Needed
see pp 1-4, 8

ERRORED SEQUENCES

OK 10 <210> SEQ ID NO: 1
11 <211> LENGTH: 1306
12 <212> TYPE: DNA
13 <213> ORGANISM: Mycoplasma gallisepticum
14 <223> OTHER INFORMATION: TTM-1 gene

E--> 15 <400> SEQUENCE: 1

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17 aatcctaaat aaataagccg ttaaatattac taaaaaatta aaaaaatggt ttttcttatc 120
18 aaccaaaatt ctctagtaat aaacgcttat ttattttttat ttttagtcat cttttaagat 180
19 ataaatatat cttaatatct tatgaataag aaaagaatca tcttaaagac tattagtttg 240
20 ttaggtacaa catcctttct tagcattggg atttctagct gtatgtctat tactaaaaaa 300
21 gatgcaaacc caaataatgg ccaaacccaa ttagaagcag cgcgatgga gttaacagat 360
22 ctaatcaatg ctaaagcgat gacattagct tcaactacaag actatgccaa gattgaagct 420
23 agttttatcat ctgcttatag tgaagctgaa acagttaaca ataaccttaa tgcaacatta 480
24 gaacaactaa aatgggctaa aactaattta gaatcagcca tcaaccaagc taatacggat 540
25 aaaacgactt ttgataatga acacccaaat ttagttgaag catacaaagc actaaaaacc 600
26 acttttagaac aacgtgctac taaccttgaa ggtttgtcat caactgctta taatcaaatt 660
27 cgcaataatt tagtggatct atacaataaa gctagtagtt taataactaa aacactagat 720
28 ccactaaatg ggggaacgct ttagattctt aatgagatta ctacagttaa tcggaatatt 780
29 aataatacgt tatcaactat taatgaacaa aagactaatg ctgatgcatt atctaatagt 840
30 tttattaaaa aagtgattca aaataatgaa caaagttttg tagggacttt tacaaacgct 900
31 aatgtttcaac cttcaaacta cagttttgtt gcttttagtg ctgatgtaac acccgtcaat 960
32 tataaatatg caagaaggac cgtttggaat ggtgatgaac cttcaagtag aattcttgca 1020
33 aacacgaata gtatcacaga tgtttcttggt atttatagtt tagctggaac aaacacgaag 1080
34 taccaatatta gtttttagcaa ctatggtcca tcaactgggt atttatattt cccttataag 1140
35 ttggttaaag cagctgatgc taataacggt ggattacaat acaaattaaa taatggaaat 1200
36 gttcaacaag ttgagtttgc cacttcaact agtgcaata atactacagc taatccaact 1260
37 ccagcagttg atgagattaa agttgctaaa atcgttttat caggtt 1306

38 <210> SEQ ID NO: 2

39 <211> LENGTH: 3189

40 <212> TYPE: DNA

41 <213> ORGANISM: Mycoplasma gallisepticum

p.2

<220> insert this mandatory
numeric
identifier
whenever
<221>, <222>
or <223>
is shown

FYI: Per
1.823 of
Sequence Rules,
"a fixed-width
font should be
used exclusively
throughout the

'Sequence Listing'
do not vary the
font in the file 10/1/02

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(2207) insert

42 <223> OTHER INFORMATION: mgc3 gene

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45	gctcttggtt	ctgcaagctt	tggctttaag	caatcagata	agagtaacga	taacacgcaa	120
46	ttagttaatc	aagcaagaac	gctagatgct	aattctgtta	gacttgcagg	tcttggacaa	180
47	aatggttcgt	tgttcaatac	agttcttaga	gatgttgatg	ataactttat	aacagcagct	240
48	aatggaacaa	ttatcaaatt	agatagtttt	actaaaccaa	tatatgggtt	agatctaaag	300
49	gatgattgtg	gtggatacaa	agtaaaacaa	atagtttcag	attacacaa	tagcagaaat	360
50	agatttgatc	aaagacaaac	aagagcatat	tatgctctgt	tgggttaatga	tgaagctaac	420
51	gttcatttaa	aaagaattaa	tactaactca	aatagaattg	gtaatagaaa	caacaattct	480
52	aagtttgtaa	ttggtggtgt	tgataatcca	gctcacgtaa	ttagattttac	tgatgatggg	540
53	actaaattta	attttacaaa	ccaaactcaa	ggtgaaattg	ttaatgactt	catttttagat	600
54	gcgccaatct	tacctaaaga	tttacaccca	gattgggtata	acttatacat	tcaaagaaa	660
57	atcttaccac	atgacgtcaa	cactgcagtt	gttccttggc	cagtaggtag	agttagtggg	720
58	acaaatgctg	atgatgggat	gtttgattgt	gggaatgggc	aaataactaa	tacagatcct	780
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60	atgcctgggtg	caaacaatag	atacgattct	caattgaatg	tcaagcatag	aattaaaaca	900
61	tctttccaat	tagatgaaaa	atttgtttat	ccagaatgga	ctggttctga	agagaataaa	960
62	aatattacaa	gattagctac	tggaagtttg	ccaagcaacg	aaagatattg	gattccttgac	1020
63	atacccgga	ctccacaagt	tactttaaaa	gaagattcag	ttaacgtatt	ttcaagacta	1080
64	tacttaaaact	cagttaattc	tttatcattc	attggtgata	gtattttatat	ttttggtacc	1140
65	tctgaattac	catcattatg	gtactattca	ttcccaacta	gattatctga	tctaaccgct	1200
66	ttgaatcaag	ttaaaacaga	tgatattgaa	gcttcaagca	ctgataacgg	tacaacaaca	1260
67	aacggaacaa	cgacaacagc	tgatacatct	agtggttcaa	caggtgctgg	aacaggaaa	1320
68	actactaaca	cttctcaaac	agtttcta	cctactttaa	atacttatcg	tagttttgga	1380
69	attgatagta	aaaccaattc	tgcaaacaaa	atagatgaaa	ctaattgggc	agatcctaac	1440
70	gttattgaag	caagaatata	tgctgaatac	agatttaggt	ttcaaaatga	aattccaata	1500
71	actaatgcag	gaaactttat	ccgaaacaca	attggtgggt	ttggttttac	ttcaacaggt	1560
72	tcaagagtag	ttttaagagc	ttcttataac	ggtgatcaac	gtccaactgg	aaacttccaa	1620
73	cctttcttat	acgtatttgg	ttatttagga	taccaacaaa	ctagaacagg	aactttctgg	1680
74	tacggaacat	ataagctttt	aaacaacagc	ccttacgacg	tattagattc	tccaagagta	1740
75	ggtactgaaa	ccaatcaatt	tagaagaact	tcattaacat	accctgttat	gggtggatat	1800
76	ctaactgaag	aaggtgctag	aagtttctct	aatactccat	atataagagc	acaaggtgac	1860
77	acaccagaaa	gccgaagcat	cttccaatct	ggctattctg	ataatactta	tgagtacatt	1920
78	caatcagttt	taggatttga	tggaattaga	aataacttaa	atgttgggg	ttaaagcatca	1980
79	agcttcttaa	actcaaatag	accaaatacca	aacggtctag	aaatgattgc	tgcaacaaca	2040
80	tacttaagat	cacaaattgg	attagctaga	acatctggat	taccaaacca	acaaccattc	2100
81	ggaacaactc	accaagttat	ttcagtatca	cctggtgatc	agttctcatc	aattaagaat	2160
82	attagaacaa	tcttccctgg	taaccagtta	tggtacttct	tattcacaaa	tgaaaataat	2220
83	aaatctagt	tttatacatt	aagattagct	gactcaagta	accctgatgc	gtcaagctca	2280
84	ttcagtccaa	caagttaaat	tgacgtta	gaaattgggt	taactttacc	tttattagac	2340
85	aattcattct	atacagtaaa	tgctgctgg	aatgttgc	tggtctcatc	aaaccctgg	2400
86	tctcctggat	catatactgc	tgtaaatcaga	tttaatcaga	acttatctga	tattgctttt	2460
87	gaaggttctg	gtgctaagta	tacatctgat	ttctggggaa	caatccaatt	caaaccgat	2520
88	gagtacttaa	ttcaaaatgg	gttcactagt	caagtggcta	gaaacttcgt	tacaaaccaa	2580
89	agcttcttaa	acagtttagt	tgacttca	cctgcta	ctggtactaa	ctaccgtgta	2640
90	gtggttgatc	ctgatggtaa	tttaacaaac	caaaacctac	ctctaaaagt	tcagatccaa	2700
91	tacttagatg	gtaagtatta	tgatgctaaa	ttaaagaaca	ataatttagt	aacattctct	2760
92	tataacaact	ttggcgctt	accttcatgg	gtagtgcta	cagcaattgg	tagtacatta	2820

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DATE: 10/01/2002

TIME: 16:12:31

Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

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93 ggtattcttg caattatgat catcttagga ttagctatcg gtattccttt aagagctcaa 2880
94 agaaaattac aagacaaagg gttcaaaaca acattcaaaa aagttgatac cttgactgct 2940
95 gctgttggtt cagtttataa gaagattatt acccaaactg ctaacgtaa gaaaaaacct 3000
96 gctgctttag gtgctggtaa atctggtgat aagaaacctg ctgctgctgc taaacctgct 3060
97 gctccagcta aaccatctgc accaaaagct agctcaccag ctaaaccaac tgggcctaaa 3120
98 tctggtgcgc ctacaaaacc aactgctcct aagccagctg ctccaaaacc aaccgctccc 3180
99 aaagaataa 3189

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E--> 100 <210> SEQ ID NO: 12

101 <211> LENGTH: 456

102 <212> TYPE: PRT

103 <213> ORGANISM: Mycoplasma gallisepticum

104 <223> OTHER INFORMATION: TTM-1 portion of pNZ40K-S

E--> 105 <400> SEQUENCE: 12

106 Met His Tyr Phe Arg Arg Asn Cys Ile Phe Phe Leu Ile Val Ile Leu

107 1 5 10 15

E--> 108 Tyr Gly Thr Asn Ser Ser Pro Ser Thr Gln Asn Val Thr Ser Arg (GIU) invalid

109 20 25 30

112 Val Val Ser Ser Val Gln Leu Ser Glu Glu Glu Ser Thr Phe Tyr Leu

113 35 40 45

114 Cys Pro Pro Pro Val Gly Ser Thr Val Ile Arg Leu Glu Phe Gly Cys

115 50 55 60

116 Met Ser Ile Thr Lys Lys Asp Ala Asn Pro Asn Asn Gly Gln Thr Gln

117 65 70 75 80

118 Leu Glu Ala Ala Arg Met Glu Leu Thr Asp Leu Ile Asn Ala Lys Ala

119 85 90 95

120 Met Thr Leu Ala Ser Leu Gln Asp Tyr Ala Lys Ile Glu Ala Ser Leu

121 100 105 110

122 Ser Ser Ala Tyr Ser Glu Ala Glu Thr Val Asn Asn Asn Leu Asn Ala

123 115 120 125

124 Thr Leu Glu Gln Leu Lys Met Ala Lys Thr Asn Leu Glu Ser Ala Ile

125 130 135 140

126 Asn Gln Ala Asn Thr Asp Lys Thr Thr Phe Asp Asn Glu His Pro Asn

127 145 150 155 160

128 Leu Val Glu Ala Tyr Lys Ala Leu Lys Thr Thr Leu Glu Gln Arg Ala

129 165 170 175

130 Thr Asn Leu Glu Gly Leu Ser Ser Thr Ala Tyr Asn Gln Ile Arg Asn

131 180 185 190

132 Asn Leu Val Asp Leu Tyr Asn Lys Ala Ser Ser Leu Ile Thr Lys Thr

133 195 200 205

134 Leu Asp Pro Leu Asn Gly Gly Thr Leu Leu Asp Ser Asn Glu Ile Thr

135 210 215 220

136 Thr Val Asn Arg Asn Ile Asn Asn Thr Leu Ser Thr Ile Asn Glu Gln

137 225 230 235 240

138 Lys Thr Asn Ala Asp Ala Leu Ser Asn Ser Phe Ile Lys Lys Val Ile

139 245 250 255

140 Gln Asn Asn Glu Gln Ser Phe Val Gly Thr Phe Thr Asn Ala Asn Val

141 260 265 270

142 Gln Pro Ser Asn Tyr Ser Phe Val Ala Phe Ser Ala Asp Val Thr Pro

143 275 280 285



<220> insert

Segs
3 through 11
missing
(see p. 8)

do you
mean Glu?

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Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

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144 Val Asn Tyr Lys Tyr Ala Arg Arg Thr Val Trp Asn Gly Asp Glu Pro
145      290      295      300
146 Ser Ser Arg Ile Leu Ala Asn Thr Asn Ser Ile Thr Asp Val Ser Trp
147 305      310      315      320
148 Ile Tyr Ser Leu Ala Gly Thr Asn Thr Lys Tyr Gln Phe Ser Phe Ser
149      325      330      335
150 Asn Tyr Gly Pro Ser Thr Gly Tyr Leu Tyr Phe Pro Tyr Lys Leu Val
151      340      345      350
152 Lys Ala Ala Asp Ala Asn Asn Val Gly Leu Gln Tyr Lys Leu Asn Asn
153      355      360      365
154 Gly Asn Val Gln Gln Val Glu Phe Ala Thr Ser Thr Ser Ala Asn Asn
155      370      375      380
156 Thr Thr Ala Asn Pro Thr Pro Ala Val Asp Glu Ile Lys Val Ala Lys
157 385      390      395      400
158 Ile Val Leu Ser Gly Leu Arg Phe Gly Gln Asn Thr Ile Glu Leu Ser
159      405      410      415
160 Val Pro Thr Gly Glu Gly Asn Met Asn Lys Val Ala Pro Met Ile Gly
161      420      425      430
162 Asn Ile Tyr Leu Ser Ser Asn Glu Asn Asn Ala Asp Lys Ile Pro Gly
163      435      440      445
164 Tyr Arg Arg Pro Gly Thr Phe Leu
165      450      455

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E--> 168 <210> SEQ ID NO: 25

169 <211> LENGTH: 1224

170 <212> TYPE: PRT

171 <213> ORGANISM: Mycoplasma gallisepticum

172 <223> OTHER INFORMATION: MGC3 encoded by mgc3 gene

E--> 173 <400> SEQUENCE: 25

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174 Met Asn Ile Ser Lys Lys Leu Lys Ser Tyr Thr Leu Ile Gly Gly Leu
175 1      5      10      15
176 Ala Val Phe Gly Ala Leu Gly Ser Ala Ser Phe Gly Phe Lys Gln Ser
177      20      25      30
178 Asp Lys Ser Asn Asp Asn Thr Gln Leu Val Asn Gln Ala Arg Thr Leu
179      35      40      45
180 Asp Ala Asn Ser Val Arg Leu Ala Gly Leu Gly Gln Asn Gly Ser Leu
181      50      55      60
182 Phe Asn Thr Val Leu Arg Asp Val Asp Asp Asn Phe Ile Thr Ala Ala
183 65      70      75      80
184 Asn Gly Thr Ile Ile Lys Leu Asp Ser Phe Thr Lys Pro Leu Tyr Gly
185      85      90      95
186 Leu Asp Leu Ser Asp Asp Cys Gly Gly Tyr Lys Val Lys Gln Ile Val
187      100      105      110
188 Ser Asp Tyr Thr Thr Ser Arg Asn Arg Phe Asp Gln Arg Gln Thr Arg
189      115      120      125
190 Ala Tyr Tyr Ala Leu Leu Val Asn Asp Glu Ala Asn Val His Leu Lys
191      130      135      140
192 Arg Ile Asn Thr Asn Ser Asn Arg Ile Gly Asn Arg Asn Asn Asn Ser
193 145      150      155      160
194 Lys Phe Val Ile Gly Gly Val Asp Asn Pro Ala His Val Ile Arg Phe

```

→ Segs 13 through 24
missing
(see p. 8)

→ 12207 insert

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PATENT APPLICATION: US/09/901,572

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Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

195				165				170				175
196	Thr	Asp	Asp	Gly	Thr	Lys	Phe	Asn	Phe	Thr	Asn	Gln
197				180				185				190
198	Ile	Val	Asn	Asp	Phe	Ile	Leu	Asp	Ala	Pro	Ile	Leu
199			195					200				205
200	His	Pro	Asp	Trp	Tyr	Asn	Leu	Tyr	Ile	Gln	Arg	Lys
201		210				215					220	
202	Asp	Val	Asn	Thr	Ala	Val	Val	Pro	Trp	Pro	Val	Gly
203	225					230				235		240
204	Thr	Asn	Ala	Asp	Asp	Gly	Met	Phe	Asp	Cys	Gly	Asn
205				245					250			255
206	Asn	Thr	Asp	Pro	Ile	Ala	Gln	Thr	Lys	Thr	Thr	Thr
207			260						265			270
208	Pro	Ser	Thr	Phe	Asn	Ser	Gly	Ala	Met	Pro	Gly	Ala
209			275					280				285
210	Asp	Ser	Gln	Leu	Asn	Val	Lys	His	Arg	Ile	Lys	Thr
211		290					295				300	
212	Asp	Glu	Arg	Ile	Asn	Thr	Asn	Ser	Asn	Arg	Ile	Gly
213	305					310				315		320
214	Asn	Ser	Lys	Phe	Val	Ile	Gly	Gly	Val	Asp	Asn	Pro
215				325					330			335
216	Arg	Phe	Thr	Asp	Asp	Gly	Thr	Lys	Phe	Asn	Phe	Thr
217			340					345				350
218	Gly	Glu	Ile	Val	Asn	Asp	Phe	Ile	Leu	Asp	Ala	Pro
219			355				360				365	
220	Asp	Leu	His	Pro	Asp	Trp	Tyr	Asn	Leu	Tyr	Ile	Gln
221		370					375				380	
224	Pro	Asn	Asp	Val	Asn	Thr	Ala	Val	Val	Pro	Trp	Pro
225	385					390				395		400
226	Ser	Gly	Thr	Asn	Ala	Asp	Asp	Gly	Met	Phe	Asp	Cys
227				405					410			415
228	Ile	Thr	Asn	Thr	Asp	Pro	Ile	Ala	Gln	Thr	Lys	Thr
229			420						425			430
230	Gln	Asn	Pro	Ser	Thr	Phe	Asn	Ser	Gly	Ala	Met	Pro
231			435				440				445	
232	Arg	Tyr	Asp	Ser	Gln	Leu	Asn	Val	Lys	His	Arg	Ile
233		450					455				460	
234	Gln	Leu	Asp	Glu	Lys	Phe	Val	Tyr	Pro	Glu	Trp	Thr
235	465					470				475		480
236	Asn	Lys	Asn	Ile	Thr	Arg	Leu	Ala	Thr	Gly	Ser	Leu
237				485					490			495
238	Arg	Tyr	Trp	Ile	Leu	Asp	Ile	Pro	Gly	Thr	Pro	Gln
239			500					505				510
240	Glu	Asp	Ser	Val	Asn	Val	Phe	Ser	Arg	Leu	Tyr	Leu
241			515					520				525
242	Ser	Leu	Ser	Phe	Ile	Gly	Asp	Ser	Ile	Tyr	Ile	Phe
243		530					535				540	
244	Leu	Pro	Ser	Leu	Trp	Tyr	Tyr	Ser	Phe	Pro	Thr	Arg
245	545					550				555		560

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Input Set : A:\EP.txt

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246 Thr Ala Leu Asn Gln Val Lys Thr Asp Asp Ile Glu Ala Ser Ser Thr
247          565          570          575
248 Asp Asn Gly Thr Thr Thr Asn Gly Thr Thr Thr Ala Asp Thr Ser
249          580          585          590
250 Ser Gly Ser Thr Gly Ala Gly Thr Gly Asn Thr Thr Asn Thr Ser Gln
251          595          600          605
252 Thr Val Ser Asn Pro Thr Leu Asn Thr Tyr Arg Ser Phe Gly Ile Asp
253          610          615          620
254 Ser Lys Pro Thr Ser Ala Asn Lys Ile Asp Glu Thr Asn Trp Ala Asp
255 625          630          635          640
256 Pro Asn Val Ile Glu Ala Arg Ile Tyr Ala Glu Tyr Arg Leu Gly Ile
257          645          650          655
258 Gln Asn Glu Ile Pro Ile Thr Asn Ala Gly Asn Phe Ile Arg Asn Thr
259          660          665          670
260 Ile Gly Gly Val Gly Phe Thr Ser Thr Gly Ser Arg Val Val Leu Arg
261          675          680          685
262 Ala Ser Tyr Asn Gly Asp Gln Arg Pro Thr Gly Asn Phe Gln Pro Phe
263          690          695          700
264 Leu Tyr Val Phe Gly Tyr Leu Gly Tyr Gln Gln Thr Arg Thr Gly Thr
265 705          710          715          720
266 Phe Trp Tyr Gly Thr Tyr Lys Leu Leu Asn Ser Pro Tyr Asp Val
267          725          730          735
268 Leu Asp Ser Pro Arg Val Gly Thr Glu Thr Asn Gln Phe Arg Arg Thr
269          740          745          750
270 Ser Leu Thr Tyr Pro Val Met Gly Gly Tyr Leu Thr Glu Glu Gly Ala
271          755          760          765
272 Arg Ser Phe Ser Asn Thr Pro Tyr Ile Arg Ala Gln Gly Asp Thr Pro
273          770          775          780
274 Glu Ser Arg Ser Ile Phe Gln Ser Gly Tyr Ser Asp Asn Thr Tyr Glu
275 785          790          795          800
276 Tyr Ile Gln Ser Val Leu Gly Phe Asp Gly Ile Arg Asn Asn Leu Asn
277          805          810          815
280 Val Gly Val Lys Ala Ser Ser Phe Leu Asn Ser Asn Arg Pro Asn Pro
281          820          825          830
282 Asn Gly Leu Glu Met Ile Ala Ala Thr Thr Tyr Leu Arg Ser Gln Ile
283          835          840          845
284 Gly Leu Ala Arg Thr Ser Gly Leu Pro Asn Gln Gln Pro Phe Gly Thr
285          850          855          860
286 Thr His Gln Val Ile Ser Val Ser Pro Gly Asp Gln Phe Ser Ser Ile
287 865          870          875          880
288 Lys Asn Ile Arg Thr Ile Phe Pro Gly Asn Gln Leu Trp Tyr Phe Leu
289          885          890          895
290 Phe Thr Asn Glu Asn Asn Lys Ser Ser Val Tyr Thr Leu Arg Leu Ala
291          900          905          910
292 Asp Ser Ser Asn Pro Asp Ala Ser Ser Ser Phe Ser Pro Thr Ser Leu
293          915          920          925
294 Ile Asp Val Asn Glu Ile Gly Val Ile Leu Pro Leu Leu Asp Asn Ser
295          930          935          940
296 Phe Tyr Thr Val Asn Ala Ala Gly Asn Val Ala Leu Phe Ser Ser Asn

```

RAW SEQUENCE LISTING

DATE: 10/01/2002

PATENT APPLICATION: US/09/901,572

TIME: 16:12:31

Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

```

297 945          950          955          960
298 Pro Gly Ser Pro Gly Ser Tyr Thr Ala Val Asn Thr Phe Asn Gln Asn
299          965          970          975
300 Leu Ser Asp Ile Ala Phe Glu Gly Ser Gly Ala Lys Tyr Thr Ser Asp
301          980          985          990
302 Phe Trp Gly Thr Ile Gln Phe Lys Pro Asp Glu Tyr Leu Ile Gln Asn
303          995          1000          1005
304 Gly Phe Thr Ser Gln Val Ala Arg Asn Phe Val Thr Asn Gln Ser Phe
305          1010          1015          1020
306 Leu Asn Ser Leu Val Asp Phe Thr Pro Ala Asn Ala Gly Thr Asn Tyr
307 1025          1030          1035          1040
308 Arg Val Val Val Asp Pro Asp Gly Asn Leu Thr Asn Gln Asn Leu Pro
309          1045          1050          1055
310 Leu Lys Val Gln Ile Gln Tyr Leu Asp Gly Lys Tyr Tyr Asp Ala Lys
311          1060          1065          1070
312 Leu Lys Asn Asn Asn Leu Val Thr Phe Ser Tyr Asn Asn Phe Gly Ala
313          1075          1080          1085
314 Leu Pro Ser Trp Val Val Pro Thr Ala Ile Gly Ser Thr Leu Gly Ile
315          1090          1095          1100
316 Leu Ala Ile Met Ile Ile Leu Gly Leu Ala Ile Gly Ile Pro Leu Arg
317 1105          1110          1115          1120
318 Ala Gln Arg Lys Leu Gln Asp Lys Gly Phe Lys Thr Thr Phe Lys Lys
319          1125          1130          1135
320 Val Asp Thr Leu Thr Ala Ala Val Gly Ser Val Tyr Lys Lys Ile Ile
321          1140          1145          1150
322 Thr Gln Thr Ala Asn Val Lys Lys Lys Pro Ala Ala Leu Gly Ala Gly
323          1155          1160          1165
324 Lys Ser Gly Asp Lys Lys Pro Ala Ala Ala Ala Lys Pro Ala Ala Pro
325          1170          1175          1180
326 Ala Lys Pro Ser Ala Pro Lys Ala Ser Ser Pro Ala Lys Pro Thr Gly
327 1185          1190          1195          1200
328 Pro Lys Ser Gly Ala Pro Thr Lys Pro Thr Ala Pro Lys Pro Ala Ala
329          1205          1210          1215
330 Pro Lys Pro Thr Ala Pro Lys Glu
331          1220

```


RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/901,572

DATE: 10/01/2002
TIME: 16:12:32

Input Set : A:\EP.txt
Output Set: N:\CRF4\10012002\I901572.raw

Skipped Sequences(NEW RULES):

Sequence(s)___missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

<400> sequence id number

000

Seq#:3,4,5,6,7,8,9,10,11,13,14,15,16,17,18,19,20,21,22,23,24

VERIFICATION SUMMARY

DATE: 10/01/2002

PATENT APPLICATION: US/09/901,572

TIME: 16:12:32

Input Set : A:\EP.txt

Output Set: N:\CRF4\10012002\I901572.raw

L:4 M:283 W: Missing Blank Line separator, <120> field identifier
L:5 M:283 W: Missing Blank Line separator, <130> field identifier
L:6 M:283 W: Missing Blank Line separator, <140> field identifier
L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:8 M:212 E: (34) Invalid or duplicate Sequence ID Number, Number Of Sequences Unknown
L:8 M:283 W: Missing Blank Line separator, <160> field identifier
L:10 M:283 W: Missing Blank Line separator, <210> field identifier
L:15 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:1
L:15 M:283 W: Missing Blank Line separator, <400> field identifier
L:43 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:2
L:43 M:283 W: Missing Blank Line separator, <400> field identifier
L:100 M:216 E: (34) Seq.#s missing, SEQ ID NOS: 3 thru 11
L:105 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:12
L:105 M:283 W: Missing Blank Line separator, <400> field identifier
L:108 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:168 M:216 E: (34) Seq.#s missing, SEQ ID NOS: 13 thru 24
L:173 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:25
L:173 M:283 W: Missing Blank Line separator, <400> field identifier
L:8 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (0) Counted (4)